

CLAIMS

What is claimed is:

1. A method for identifying compounds that affect LDL-proteoglycan binding, comprising the steps of:
 - (a) incubating a mixture comprising (i) proteoglycan, (ii) LDL, and (iii) a candidate compound, under conditions wherein LDL binds to proteoglycan to form an LDL-proteoglycan complex in the absence of said candidate compound;
 - (b) determining any difference between the amount of LDL-proteoglycan complex present in:
 - (i) the mixture prepared in step (a), and
 - (ii) a assay mixture comprising said proteoglycan and said LDL in the absence of said candidate compound.
2. The method according to claim 1, further comprising the step of:
 - (c) correlating any difference determined in step (b) with said candidate compound's ability to affect LDL-proteoglycan binding.
3. The method according to claim 1 or 2, wherein the LDL of step (a) is attached to a solid support.
4. The method according to claim 1 or 2, wherein the proteoglycan of step (a) is attached to a solid support.
5. The method according to claim 1, 2 or 4, wherein the LDL of step (a) is labeled.
6. The method according to claim 3, wherein the proteoglycan of step (a) is labeled.
7. The method according to claim 6, wherein the label is biotin.

13. The compounds which affect LDL-proteoglycan binding, which do not substantially affect LDL receptor binding identified by the method according to claim 10 or 11.

14. A apo-B100 protein comprising a proteoglycan receptor⁺ mutation in Site B.

15. The apo-B100 protein according to claim 14, which is purified.

16. The apo-B100 protein according to claim 14, which is synthesized by recombinant DNA expression or chemical synthesis.

17. The apo-B100 protein according to claim 14, wherein the amino acid sequence from position 3358 to 3359 is selected from the group consisting of:

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Glu**₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Asp**₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Ala**₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Thr**₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Ser**₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Gln**₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-**Glu**₃₃₆₂-Lys₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-**Asp**₃₃₆₂-Lys₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-Lys₃₃₆₃-**Glu**₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-Lys₃₃₆₃-**Asp**₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-**Glu**₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-Lys₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-**Asp**₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-Lys₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-----Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-----Lys₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-Lys₃₃₆₃-----Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Glu**-Lys₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-Lys₃₃₆₃-**Glu**-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇,
Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Asp**-Lys₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇, and
Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-Lys₃₃₆₃-**Asp**-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇.

18. The apo-B100 protein according to claim 14, wherein said mutation in Site B is the K3363E mutation, and the amino acid sequence from position 3358 to 3359 is:

Thr₃₃₅₈-Arg₃₃₅₉-Leu₃₃₆₀-Thr₃₃₆₁-Arg₃₃₆₂-**Glu**₃₃₆₃-Arg₃₃₆₄-Gly₃₃₆₅-Leu₃₃₆₆-Lys₃₃₆₇.

19. A polypeptide comprising the amino acid sequence of Site B in the apo-B100 protein according to any one of claims 14 to 18, wherein said Site B is flanked on at least one side by a contiguous sequence of at least 10 amino acids which is directly adjacent to Site B in the wild-type human apo-B100 sequence.

20. An LDL particle comprising an apo-B100 protein according to any one of claims 14 to 18.

21. An LDL particle comprising a polypeptide according to claim 19.

22. An antibody composition which binds to an antigenic determinant in an apo-B100 protein according to any one of claims 14 to 18, wherein said antigenic determinant is not present in the wild-type human apo-B100 protein.

23. A polynucleotide encoding an apo-B100 protein according to any one of claims 14 to 18.

24. The polynucleotide according to claim 23, wherein the polynucleotide is present in a 95-kb apo-B P1 plasmid p158.

25. The polynucleotide according to claim 24, wherein said mutation in Site B is the K3363E mutation.

